EXHIBIT A

Claim Amendments: Pending Claims

- 1. An ApoA-I agonist compound comprising:
- (i) a 15 to 26- residue peptide or peptide analogue according to formula (I) which forms an amphipathic α -helix in the presence of lipids and exhibits at least about 38% LCAT activation activity as compared with human ApoA-I wherein one or two helical turns are deleted from formula (I), wherein a helical turn consists of 3 to 4 consecutive residues selected from residues X_1 to X_{23} of formula (I):

 $Z_{1}-X_{1}-X_{2}-X_{3}-X_{4}-X_{5}-X_{6}-X_{7}-X_{8}-X_{9}-X_{10}-X_{11}-X_{12}-X_{13}-X_{14}-X_{15}-X_{16}-X_{17}-X_{18}-X_{19}-X_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-Z_{20}-X_{21}-X_{22}-X_{23}-X_{22}-X_{23}-X_{$

or a pharmaceutically acceptable salt thereof, wherein:

- X₁ is Pro (P), Ala (A), Gly (G), Gln (Q), Asn (N), Asp (D) or D-Pro (p);
- X₂ is an aliphatic residue;
- X_3 is a Leu (L) or Phe (F);
- X_4 is Glu (E)
- X₅ is an aliphatic residue;
- X_6 is Leu (L) or Phe (F);
- X_7 is Glu (E) or Leu (L);
- X_8 is Asn (N) or Gln (Q);
- X_9 is Leu (L);
- X_{10} is Leu (L), Trp (W) or Gly (G);
- X_{11} is an acidic residue;
- X_{12} is Arg (R);
- X_{13} is Leu (L) or Gly (G);
- X_{14} is Leu (L), Phe (F) or Gly (G);
- X_{15} is Asp (D);
- X_{16} is Ala (A);
- X_{17} is Leu (L);
- X_{18} is Asn (N) or Gln (Q);
- X₁₉ is a basic residue;
- X₂₀ is a basic residue;
- X_{21} is Leu (L);

 X_{21} is Leu (L);

 X_{22} is a basic residue;

X₂₃ is absent or a basic residue;

 Z_1 is H_2N_- ;

 Z_2 is -C (O) NRR or -C (O) OR;

each R is independently -H, (C_1-C_6) alkyl, (C_1-C_6) alkenyl, (C_1-C_6) alkynyl, (C_5-C_{20}) aryl, (C_6-C_{26}) alkaryl, 5-20 membered heteroaryl or 6-26 membered alkheteroaryl or a 1 to 7-residue peptide or peptide analogue in which one more bonds between residues 1-7 are independently a substituted amide, an isostere of an amide or an amide mimetic; and

each "-" between residues X_1 to X_{23} and between residues of the peptide to Z_2 independently designates an amide linkage, a substituted amide linkage, an isostere of an amide or an amide mimetic; or

an N- terminally blocked form, a C-terminally blocked form, or an N- and C-terminally blocked form of formula (I).

- 56. The 15 to 26-residue peptide or deleted peptide analogue of Claim 1, in which one helical turn is deleted.
- 57. The 15 to 26-residue peptide or peptide analogue of Claim 1, in which three, four, six, seven or eight residues X_1 , X_2 , X_3 , X_4 , X_5 , X_6 , X_7 , X_8 , X_9 , X_{10} , X_{11} , X_{12} , X_{13} , X_{14} , X_{15} , X_{16} , X_{17} , X_{18} , X_{19} , X_{20} , X_{21} and X_{22} are deleted.
- 58. The 15 to 26-residue peptide or peptide analogue of Claim 57, in which 3 consecutive residues are deleted.
- 59. The 15 to 26-residue peptide or peptide analogue of Claim 57, in which 4 consecutive residues are deleted.
- 60. The 15 to 26-residue peptide or peptide analogue of Claim 57, in which two non-contiguous sets of 3 consecutive residues are deleted.
- 61. The 15 to 26-residue peptide or peptide analogue of Claim 57, in which two non-contiguous sets of 4 consecutive residues are deleted.

- 62. The 15 to 26-residue peptide or peptide analogue of Claim 57, in which one set of 3 consecutive residues and one set of 4 consecutive residues are deleted.
- 63. The 15 to 26-residue peptide or peptide analogue of Claim 57, in which 6, 7 or 8 consecutive residues are deleted.
- 67. The 15 to 26-residue peptide or peptide analogue of Claim 1 in which:

the "-" between residues designates -C (O) NH-;

 Z_1 is H_2N_- ; and

Z₂ is -C (O) OH or a salt thereof.

- 68. The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobic moment, $\langle \mu_H \rangle$, is 0.45 to 0.65.
- 69. The 15 to 26-residue peptide or peptide analogue of Claim 68, in which the mean hydrophobic moment, $\langle \mu_H \rangle$, is 0.50 to 0.60.
- 70. The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobicity, $\langle H_o \rangle$, is -0.050 to -0.070.
- 71. The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobicity, $\langle H_o \rangle$, is -0.030 to -0.055.
- 72. The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobicity of the hydrophobic face, $\langle H_o^{pho} \rangle$, is 0.90 to 1.20.
- 73. The 15 to 26-residue peptide or peptide analogue of Claim 72, in which the mean hydrophobicity of the hydrophobic face, $\langle H_o^{pho} \rangle$, is 0.94 to 1.10.
- 74. The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the pho angle is 160° to 220°.
- 75. The 15 to 26-residue peptide or peptide analogue of Claim 74, in which the pho angle is 180° to 200°.
- 79. A pharmaceutical composition comprising an ApoA-I agonist compound and a pharmaceutically acceptable carrier, excipient or diluent, wherein the ApoA-I agonist

compound is a 15 to 26-residue peptide or peptide analogue according to Claim 1 or 57.

- 82. The pharmaceutical composition of Claim 79 which is a lyophilized powder.
- 83. The pharmaceutical composition of Claim 79 which is a solution.
- 84. The N-terminally blocked form of the 15 to 26-residue peptide or peptide analogue of Claim 1.
- 85. The 15 to 26-residue peptide or peptide analogue of Claim 84 in which the N-terminally blocking group is selected from the group consisting of acetyl, formyl and dansyl.
- 86. The C-terminally blocked form of the 15 to 26-residue peptide or peptide analogue of Claim 1.
- 87. The 15 to 26-residue peptide or peptide analogue of Claim 86 in which the C-terminally blocking group is methyl.
- 88. The N-terminally and C-terminally blocked form of the 15 to 26-residue peptide or peptide analogue of Claim 1.